## **AMENDMENTS TO THE CLAIMS**

This listing of claims will replace all prior versions and listings of claims in the abovereferenced application.

## **Listing of Claims:**

- 1. (Currently amended) A protein of the following as defined in (a) or (b):
  - (a) a protein comprising an the amino acid sequence of SEQ ID NO: 2; and
- (b) a protein comprising an the amino acid sequence of SEQ ID NO: 2 with one or several amino acids deleted, replaced, or added, and having an activity of binding rabconnectin-3 and a GDP/GTP exchange protein.
- 2. (Currently amended) A <u>The</u> protein according to of claim 1, which has the amino acid sequence of SEQ ID NO: 2.
- 3. (Currently amended) A polynucleotide for encoding that encodes the a protein as defined in claim 1 or 2.
- 4. (Currently amended) A The polynucleotide according to of claim 3, comprising a nucleotide sequence of nucleotide numbers 1 to 4470 of a the nucleotide sequence of SEQ ID NO: 1.
- 5. (Currently amended) A polynucleotide of the following as defined in (a) or (b):
- (a) a polynucleotide comprising a nucleotide sequence of nucleotide numbers 1 to 4470 of a the nucleotide sequence of SEQ ID NO: 1; and
- (b) a polynucleotide which hybridizes with the <u>a</u> polynucleotide comprising a nucleotide sequence that is complementary to the nucleotide sequence of nucleotide numbers 1 to 4470 of the nucleotide sequence of SEQ ID NO: 1 under a stringent condition, and encodes a protein having an activity of binding rabconnectin-3 and a GDP/GTP exchange protein.

- 6. (Currently amended) A polynucleotide of the following as defined in (a) or (b):
- (a) a polynucleotide comprising a nucleotide sequence of nucleotide numbers 1 to 4470 of a the nucleotide sequence of SEQ ID NO: 1; and
- (b) a polynucleotide comprising a nucleotide sequence whose homology to the nucleotide sequence of nucleotide numbers 1 to 4470 of the nucleotide sequence of SEQ ID NO: 1 is 80% or higher and encoding a protein having an activity of binding rabconnectin-3 and a GDP/GTP exchange protein.
- 7. (Currently amended) A recombinant vector comprising the <u>a</u> polynucleotide as defined in any one of claims 3 to 6.
- 8. (Currently amended) A transformant obtained by transforming a host with the <u>a</u> polynucleotide as defined in any one of <del>claim</del> <u>claims</u> 3 to 6.
- 9. (Currently amended) A method of producing a protein having an activity of binding rabconnectin-3 and a GDP/GTP exchange protein, comprising:

culturing the <u>a</u> transformant as defined in claim 8 <u>in a culture</u>; and collecting, from <u>a the</u> culture, the <u>a</u> protein having the <u>an</u> activity of binding the rabconnectin-3 and the GDP/GTP exchange protein, expressed by the transformant.

## 10. (Canceled)

- 11. (Currently amended) A method of analyzing the <u>a first</u> polynucleotide as defined in any one of claims 3 to 6, comprising hybridizing a probe or a primer <u>with the first</u> <u>polynucleotide</u>, wherein the probe or primer includes <u>including</u> a <u>second</u> polynucleotide having at least 15 nucleotides complementary to the <u>first</u> polynucleotide <del>as defined in any one of claim 3</del> to 6 with a subject polynucleotide.
- 12. (Currently amended) An The analyzing method according to of claim 11, wherein the subject first polynucleotide is present in a subject tissue or a subject cell.

Page 4 of 14

- 13. (Currently amended) A method of analyzing a gene first polynucleotide encoding the a protein as defined in claim 1 or 2, comprising hybridizing a probe or a primer with the first polynucleotide, wherein the probe or primer includes including a second polynucleotide having at least 15 nucleotides complementary to the first polynucleotide as defined in any one of claim 3 to 6 with a subject polynucleotide.
- 14. (Currently amended) A The analyzing method of analyzing a gene according to claim 12 13, wherein the subject first polynucleotide is present in a subject tissue or a subject cell.
- 15. (Currently amended) A method of analyzing a gene, comprising the steps of:
  amplifying an mRNA in a subject tissue or a subject cell by an RT-PCR method with a
  primer that includes including a polynucleotide having at least 15 nucleotides complementary to
  the a polynucleotide as defined in any one of claims 3 to 6, and

measuring the polynucleotide as defined in any one of claim 3 to 6.

- 16. (Currently amended) An antisense polynucleotide which hybridizes with an mRNA encoding the a protein as defined in claim 1 or 2.
- 17. (Currently amended) A ribozyme for cutting an mRNA encoding the a protein as defined in claim 1 or 2.
- 18. (Currently amended) A double-stranded RNA for cutting an mRNA encoding the <u>a</u> protein as defined in claim 1 or 2 by RNA interference.
- 19. (Currently amended) An antibody against the a protein as defined in claim 1 or 2.
- 20. (Currently amended) A method of immunohistologically analyzing the a protein as defined in claim 1 or 2, comprising contacting the protein which uses the with an antibody as defined in claim 19.

- 21. (Currently amended) An The analyzing method according to of claim 20, further comprising wherein the analyzing method comprises analyzing localization determining the location of a the protein.
- 22. (Currently amended) An The analysis method according to of claim 20, further comprising wherein the analyzing method comprises analyzing determining the an amount of expression of a the protein.
- 23. (Currently amended) A method of screening a candidate for a material that promotes or inhibits of a material for promoting or inhibiting binding between a rabconnectin-3-binding protein which is the protein as defined in claim 1 or 2 or a heterogeneous heterogeneous homologous protein thereof, and a rabconnectine-3 rabconnectin-3, comprising the steps of:

reacting contacting the a protein as defined in claim 1 or 2 or a heterogeneous homologous protein thereof rabconnectin-3-binding protein with the rabconnectin-3 in the presence and absence of the candidate material materials, and

selecting the <u>a candidate</u> material which increases or decreases the binding <u>between the protein and rabconnectin-3</u>.

24. (Currently amended) A method of screening a candidate for a material that promotes or inhibits of a material for promoting or inhibiting binding between a Rab GDP/GTP exchange protein binding protein which is the protein as defined in claim 1 or 2 or a heterogeneous heterogeneous homologous protein thereof, and a Rab 3 GDP/GTP exchange protein, comprising the steps of:

homologous protein thereof Rab3-GDP/GTP exchange protein binding protein with the Rab3-GDP/GTP exchange protein binding protein with the Rab3-GDP/GTP exchange protein in the presence and absence of the candidate material materials, and selecting the a candidate material which increases or decreases the binding between the protein and Rab3-GDP/GTP exchange protein.

25. (New) A polynucleotide that encodes a protein as defined in claim 2.

Page 6 of 14

- 26. (New) A recombinant vector comprising a polynucleotide as defined in claim 25.
- 27. (New) A transformant obtained by transforming a host with a polynucleotide as defined in claim 25.
- 28. (New) A method of producing a protein having an activity of binding rabconnectin-3 and a GDP/GTP exchange protein, comprising:

culturing a transformant as defined in claim 27 in a culture; and collecting, from the culture, a protein having an activity of binding rabconnectin-3 and the GDP/GTP exchange protein.

- 29. (New) A method of analyzing a first polynucleotide as defined in claim 25, comprising hybridizing a probe or a primer with the first polynucleotide, wherein the probe or primer includes a second polynucleotide having at least 15 nucleotides complementary to the first polynucleotide.
- 30. (New) The analyzing method of claim 11, wherein the first polynucleotide is present in a tissue or a cell.
- 31. (New) A method comprising amplifying an mRNA in a tissue or a cell by an RT-PCR method with a primer that includes a polynucleotide having at least 15 nucleotides complementary to a polynucleotide as defined in claim 25.